



## SEQUENCE LISTING

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<120> DETECTION AND TREATMENT OF POLYCYSTIC KIDNEY DISEASE

<130> JHU1680-2

<140> US 09/904,968

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<170> PatentIn version 3.0

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 gg 53522

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 <212> PRT  
 <213> Homo sapiens

<400> 2

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 Leu Trp Leu Gly Ala Leu Ala Gly Gly Pro Gly Arg Gly Cys Gly Pro  
 20 25 30  
 Cys Glu Pro Pro Cys Leu Cys Gly Pro Ala Pro Gly Ala Ala Cys Arg

35	40	45
Val Asn Cys Ser Gly Arg 50	Gly Leu Arg Thr Leu 55	Gly Pro Ala Leu Arg 60
Ile Pro Ala Asp Ala Thr 65	Glu Leu Asp Val Ser 70	His Asn Leu Leu Arg 75 80
Ala Leu Asp Val 85	Gly Leu Leu Ala Asn Leu 90	Ser Ala Leu Ala Glu Leu 95
Asp Ile Ser 100	Asn Asn Lys Ile Ser Thr 105	Leu Glu Glu Gly Ile Phe Ala 110
Asn Leu Phe 115	Asn Leu Ser Glu Ile Asn Leu 120	Ser Gly Asn Pro Phe Glu 125
Cys Asp Cys Gly Leu Ala 130	Trp Leu Pro Gln Trp 135	Ala Glu Glu Gln Gln 140
Val Arg Val Val Gln Pro 145	Glu Ala Ala Thr Cys 150	Ala Gly Pro Gly Ser 155 160
Leu Ala Gly Gln Pro Leu 165	Leu Gly Ile Pro Leu Leu 170	Asp Ser Gly Cys 175
Gly Glu Glu Tyr Val Ala 180	Cys Leu Pro Asp Asn Ser 185	Ser Ser Gly Thr Val 190
Ala Ala Val Ser Phe Ser 195	Ala Ala His Glu Gly Leu 200	Leu Gln Pro Glu 205
Ala Cys Ser Ala Phe Cys 210	Phe Ser Thr Gly Gln Gly 215	Leu Ala Ala Leu 220
Ser Glu Gln Gly Trp Cys 225	Leu Cys Gly Ala Ala 230	Gln Pro Ser Ser Ala 235 240
Ser Phe Ala Cys 245	Leu Ser Leu Cys Ser Gly 250	Pro Pro Ala Pro Pro Ala 255
Pro Thr Cys Arg Gly Pro 260	Thr Leu Leu Gln His Val 265	Phe Pro Ala Ser 270
Pro Gly Ala Thr Leu Val 275	Gly Pro His Gly Pro Leu 280	Ala Ser Gly Gln 285
Leu Ala Ala Phe His Ile 290	Ala Ala Pro Leu Pro Val 295	Thr Asp Thr Arg 300
Trp Asp Phe Gly Asp Gly 305	Ser Ala Glu Val Asp 310	Ala Ala Gly Pro Ala 315 320
Ala Ser His Arg Tyr Val 325	Leu Pro Gly Arg Tyr His 330	Val Thr Ala Val 335
Leu Ala Leu Gly Ala Gly 340	Ser Ala Leu Leu Gly Thr 345	Asp Val Gln Val 350
Glu Ala Ala Pro Ala Ala 355	Leu Glu Leu Val Cys Pro 360	Ser Ser Val Gln 365

Ser Asp Glu Ser Leu Asp Leu Ser Ile Gln Asn Arg Gly Gly Ser Gly  
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 Leu Glu Ala Ala Tyr Ser Ile Val Ala Leu Gly Glu Glu Pro Ala Arg  
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 Ala Val His Pro Leu Cys Pro Ser Asp Thr Glu Ile Phe Pro Gly Asn  
 405 410 415  
 Gly His Cys Tyr Arg Leu Val Val Glu Lys Ala Ala Trp Leu Gln Ala  
 420 425 430  
 Gln Glu Gln Cys Gln Ala Trp Ala Gly Ala Ala Leu Ala Met Val Asp  
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 Ser Pro Ala Val Gln Arg Phe Leu Val Ser Arg Val Thr Arg Ser Leu  
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 Asp Val Trp Ile Gly Phe Ser Thr Val Gln Gly Val Glu Val Gly Pro  
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 Pro Gly Glu Pro His Pro Ala Thr Ala Glu His Cys Val Arg Leu Gly  
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 Pro Glu Ser Arg Ser Pro Asp Asn Arg Thr Gln Leu Ala Pro Ala Cys  
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 Met Pro Gly Gly Arg Trp Cys Pro Gly Ala Asn Ile Cys Leu Pro Leu  
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 Asp Ala Ser Cys His Pro Gln Ala Cys Ala Asn Gly Cys Thr Ser Gly  
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 Pro Gly Leu Pro Gly Ala Pro Tyr Ala Leu Trp Arg Glu Phe Leu Phe  
 675 680 685



Ser Val Pro Ala Gly Pro Pro Ala Gln Tyr Ser Val Thr Leu His Gly  
 690 695 700  
 Gln Asp Val Leu Met Leu Pro Gly Asp Leu Val Gly Leu Gln His Asp  
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 Ala Gly Pro Gly Ala Leu Leu His Cys Ser Pro Ala Pro Gly His Pro  
 725 730 735  
 Gly Pro Arg Ala Pro Tyr Leu Ser Ala Asn Ala Ser Ser Trp Leu Pro  
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 His Leu Pro Ala Gln Leu Glu Gly Thr Trp Gly Cys Pro Ala Cys Ala  
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 Leu Arg Leu Leu Ala Gln Arg Glu Gln Leu Thr Val Leu Leu Gly Leu  
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 820 825 830  
 Asp Gly Arg Leu Tyr Val Pro Thr Asn Gly Ser Ala Leu Val Leu Gln  
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 Val Asp Ser Gly Ala Asn Ala Thr Ala Thr Ala Arg Trp Pro Gly Gly  
 850 855 860  
 Ser Leu Ser Ala Arg Phe Glu Asn Val Cys Pro Ala Leu Val Ala Thr  
 865 870 875 880  
 Phe Val Pro Ala Cys Pro Trp Glu Thr Asn Asp Thr Leu Phe Ser Val  
 885 890 895  
 Val Ala Leu Pro Trp Leu Ser Glu Gly Glu His Val Val Asp Val Val  
 900 905 910  
 Val Glu Asn Ser Ala Ser Arg Ala Asn Leu Ser Leu Arg Val Thr Ala  
 915 920 925  
 Glu Glu Pro Ile Cys Gly Leu Arg Ala Thr Pro Ser Pro Glu Ala Arg  
 930 935 940  
 Val Leu Gln Gly Val Leu Val Arg Tyr Ser Pro Val Val Glu Ala Gly  
 945 950 955 960  
 Ser Asp Met Val Phe Arg Trp Thr Ile Asn Asp Lys Gln Ser Leu Thr  
 965 970 975  
 Phe Gln Asn Val Val Phe Asn Val Ile Tyr Gln Ser Ala Ala Val Phe  
 980 985 990  
 Lys Leu Ser Leu Thr Ala Ser Asn His Val Ser Asn Val Thr Val Asn  
 995 1000 1005  
 Tyr Asn Val Thr Val Glu Arg Met Asn Arg Met Gln Gly Leu Gln

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Leu Thr Ala Gly Val Leu	Val Asp Ser Ala Val Glu	Val Ala Phe		
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Pro Pro Tyr Asn Glu Ser	Phe Pro Val Pro Asp Pro	Ser Val Ala		
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Gln Val Leu Val Glu His	Asn Val Thr His Thr Tyr	Ala Ala Pro		
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Tyr Thr Trp Asp Phe Gly	Asp Gly Ser Pro Val Leu	Thr Gln Ser		
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Gln Pro Ala Ala Asn His	Thr Tyr Ala Ser Arg Gly	Thr Tyr His		
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Ala Asp Val Arg Val Phe	Glu Glu Leu Arg Gly Leu	Ser Val Asp		
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1220	1225	1230		
Ala Val Gln Thr Gly Asp	Asn Ile Thr Trp Thr Phe	Asp Met Gly		
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Pro Ala Gly His Leu Ala	Arg Ser Leu His Val Leu	Val Phe Val		
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Pro Asp Ala Arg Leu Thr	Ala Tyr Val Thr Gly Asn	Pro Ala His		
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Tyr	Leu	Phe	Asp	Trp	Thr	Phe	Gly	Asp	Gly	Ser	Ser	Asn	Thr	Thr
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His	Tyr	Phe	Thr	Ser	Ile	Cys	Val	Glu	Pro	Glu	Val	Gly	Asn	Val
1370						1375						1380		
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1475						1480						1485		
Ala	Val	Gly	Arg	Gly	Arg	Pro	Ala	Ser	Tyr	Leu	Trp	Asp	Leu	Gly
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Ser Leu Thr Val Leu Glu Ala	Gly Thr Tyr His Val Gln Leu Arg
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Phe Val Glu Pro Val Gly Trp	Leu Met Val Ala Ala Ser Pro Asn
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Pro Ala Ala Val Asn Thr Ser	Val Thr Leu Ser Ala Glu Leu Ala
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Ser Trp Glu Thr Ser Glu Pro	Phe Thr Thr His Ser Phe Pro Thr
1760	1765 1770
Pro Gly Leu His Leu Val Thr	Met Thr Ala Gly Asn Pro Leu Gly
1775	1780 1785
Ser Ala Asn Ala Thr Val Glu	Val Asp Val Gln Val Pro Val Ser
1790	1795 1800
Gly Leu Ser Ile Arg Ala Ser	Glu Pro Gly Gly Ser Phe Val Ala
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Ala Gly Ser Ser Val Pro Phe	Trp Gly Gln Leu Ala Thr Gly Thr
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Asn Val Ser Trp Cys Trp Ala	Val Pro Gly Gly Ser Ser Lys Arg
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Gly Pro His Val Thr Met Val	Phe Pro Asp Ala Gly Thr Phe Ser
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Ile Arg Leu Asn Ala Ser Asn	Ala Val Ser Trp Val Ser Ala Thr
1865	1870 1875
Tyr Asn Leu Thr Ala Glu Glu	Pro Ile Val Gly Leu Val Leu Trp
1880	1885 1890
Ala Ser Ser Lys Val Val Ala	Pro Gly Gln Leu Val His Phe Gln
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Ile Leu Leu Ala Ala Gly Ser	Ala Val Thr Phe Arg Leu Gln Val
1910	1915 1920
Gly Gly Ala Asn Pro Glu Val	Leu Pro Gly Pro Arg Phe Ser His

1925	1930	1935
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Asn His Val Ser Trp Ala 1955	Gln Ala Gln Val Arg 1960	Ile Val Val Leu 1965
Glu Ala Val Ser Gly Leu 1970	Gln Val Pro Asn Cys 1975	Cys Glu Pro Gly 1980
Ile Ala Thr Gly Thr Glu 1985	Arg Asn Phe Thr Ala 1990	Arg Val Gln Arg 1995
Gly Ser Arg Val Ala Tyr 2000	Ala Trp Tyr Phe Ser 2005	Leu Gln Lys Val 2010
Gln Gly Asp Ser Leu Val 2015	Ile Leu Ser Gly Arg 2020	Asp Val Thr Tyr 2025
Thr Pro Val Ala Ala Gly 2030	Leu Leu Glu Ile Gln 2035	Val Arg Ala Phe 2040
Asn Ala Leu Gly Ser Glu 2045	Asn Arg Thr Leu Val 2050	Leu Glu Val Gln 2055
Asp Ala Val Gln Tyr Val 2060	Ala Leu Gln Ser Gly 2065	Pro Cys Phe Thr 2070
Asn Arg Ser Ala Gln Phe 2075	Glu Ala Ala Thr Ser 2080	Pro Ser Pro Arg 2085
Arg Val Ala Tyr His Trp 2090	Asp Phe Gly Asp Gly 2095	Ser Pro Gly Gln 2100
Asp Thr Asp Glu Pro Arg 2105	Ala Glu His Ser Tyr 2110	Leu Arg Pro Gly 2115
Asp Tyr Arg Val Gln Val 2120	Asn Ala Ser Asn Leu 2125	Val Ser Phe Phe 2130
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Pro Glu Val Asp Val Val 2150	Leu Pro Leu Gln Val 2155	Leu Met Arg Arg 2160
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Ser Cys Gln Arg Pro Gly 2195	Arg Pro Ala Arg Val 2200	Ala Leu Pro Gly 2205
Val Asp Val Ser Arg Pro 2210	Arg Leu Val Leu Pro 2215	Arg Leu Ala Leu 2220
Pro Val Gly His Tyr Cys 2225	Phe Val Phe Val Val 2230	Ser Phe Gly Asp 2235

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2270			2275			2280		
Pro Asn	Leu Glu	Asp Gly	Asp	Gln Thr	Pro Leu	Ser	Phe His	Trp
2285			2290			2295		
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2300			2305			2310		
Asn Phe	Gly Pro	Arg Gly	Ser	Ser Thr	Val Thr	Ile	Pro Arg	Glu
2315			2320			2325		
Arg Leu	Ala Ala	Gly Val	Glu	Tyr Thr	Phe Ser	Leu	Thr Val	Trp
2330			2335			2340		
Lys Ala	Gly Arg	Lys Glu	Glu	Ala Thr	Asn Gln	Thr	Val Leu	Ile
2345			2350			2355		
Arg Ser	Gly Arg	Val Pro	Ile	Val Ser	Leu Glu	Cys	Val Ser	Cys
2360			2365			2370		
Lys Ala	Gln Ala	Val Tyr	Glu	Val Ser	Arg Ser	Ser	Tyr Val	Tyr
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2390			2395			2400		
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2510			2515			2520		
Glu Phe	Cys Val	Tyr Lys	Gly	Ser Leu	Ser Ser	Tyr	Gly Ala	Val
2525			2530			2535		

Leu Pro	Pro Gly Phe Arg	Pro	His Phe Glu Val Gly	Leu Ala Val
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Val Val	Gln Asp Gln Leu Gly	Ala Ala Val Val	Ala	Leu Asn Arg
2555		2560		2565
Ser Leu	Ala Ile Thr Leu Pro	Glu Pro Asn Gly	Ser	Ala Thr Gly
2570		2575		2580
Leu Thr	Val Trp Leu His Gly	Leu Thr Ala Ser	Val	Leu Pro Gly
2585		2590		2595
Leu Leu	Arg Gln Ala Asp Pro	Gln His Val Ile	Glu	Tyr Ser Leu
2600		2605		2610
Ala Leu	Val Thr Val Leu Asn	Glu Tyr Glu Arg	Ala	Leu Asp Val
2615		2620		2625
Ala Ala	Glu Pro Lys His Gly	Arg Gln His Arg	Ala	Gln Ile Arg
2630		2635		2640
Lys Asn	Ile Thr Glu Thr Leu	Val Ser Leu Arg	Val	His Thr Val
2645		2650		2655
Asp Asp	Ile Gln Gln Ile Ala	Ala Ala Leu Ala	Gln	Cys Met Gly
2660		2665		2670
Pro Ser	Arg Glu Leu Val Cys	Arg Ser Cys Leu	Lys	Gln Thr Leu
2675		2680		2685
His Lys	Leu Glu Ala Met Met	Leu Ile Leu Gln	Ala	Glu Thr Thr
2690		2695		2700
Ala Gly	Thr Val Thr Pro Thr	Ala Ile Gly Asp	Ser	Ile Leu Asn
2705		2710		2715
Ile Thr	Gly Asp Leu Ile His	Leu Ala Ser Ser	Asp	Val Arg Ala
2720		2725		2730
Pro Gln	Pro Ser Glu Leu Gly	Ala Glu Ser Pro	Ser	Arg Met Val
2735		2740		2745
Ala Ser	Gln Ala Tyr Asn Leu	Thr Ser Ala Leu	Met	Arg Ile Leu
2750		2755		2760
Met Arg	Ser Arg Val Leu Asn	Glu Glu Pro Leu	Thr	Leu Ala Gly
2765		2770		2775
Glu Glu	Ile Val Ala Gln Gly	Lys Arg Ser Asp	Pro	Arg Ser Leu
2780		2785		2790
Leu Cys	Tyr Gly Gly Ala Pro	Gly Pro Gly Cys	His	Phe Ser Ile
2795		2800		2805
Pro Glu	Ala Phe Ser Gly Ala	Leu Ala Asn Leu	Ser	Asp Val Val
2810		2815		2820
Gln Leu	Ile Phe Leu Val Asp	Ser Asn Pro Phe	Pro	Phe Gly Tyr
2825		2830		2835
Ile Ser	Asn Tyr Thr Val Ser	Thr Lys Val Ala	Ser	Met Ala Phe

2840		2845		2850
Gln Thr	Gln Ala Gly Ala	Gln Ile Pro Ile Glu Arg	Leu Ala Ser	
2855		2860		2865
Glu Arg	Ala Ile Thr Val	Lys Val Pro Asn Asn Ser	Asp Trp Ala	
2870		2875		2880
Ala Arg	Gly His Arg Ser	Ser Ala Asn Ser Ala Asn	Ser Val Val	
2885		2890		2895
Val Gln	Pro Gln Ala Ser	Val Gly Ala Val Val Thr	Leu Asp Ser	
2900		2905		2910
Ser Asn	Pro Ala Ala Gly	Leu His Leu Gln Leu Asn	Tyr Thr Leu	
2915		2920		2925
Leu Asp	Gly His Tyr Leu	Ser Glu Glu Pro Glu Pro	Tyr Leu Ala	
2930		2935		2940
Val Tyr	Leu His Ser Glu	Pro Arg Pro Asn Glu His	Asn Cys Ser	
2945		2950		2955
Ala Ser	Arg Arg Ile Arg	Pro Glu Ser Leu Gln Gly	Ala Asp His	
2960		2965		2970
Arg Pro	Tyr Thr Phe Phe	Ile Ser Pro Gly Ser Arg	Asp Pro Ala	
2975		2980		2985
Gly Ser	Tyr His Leu Asn	Leu Ser Ser His Phe Arg	Trp Ser Ala	
2990		2995		3000
Leu Gln	Val Ser Val Gly	Leu Tyr Thr Ser Leu Cys	Gln Tyr Phe	
3005		3010		3015
Ser Glu	Glu Asp Met Val	Trp Arg Thr Glu Gly Leu	Leu Pro Leu	
3020		3025		3030
Glu Glu	Thr Ser Pro Arg	Gln Ala Val Cys Leu Thr	Arg His Leu	
3035		3040		3045
Thr Ala	Phe Gly Ala Ser	Leu Phe Val Pro Pro Ser	His Val Arg	
3050		3055		3060
Phe Val	Phe Pro Glu Pro	Thr Ala Asp Val Asn Tyr	Ile Val Met	
3065		3070		3075
Leu Thr	Cys Ala Val Cys	Leu Val Thr Tyr Met Val	Met Ala Ala	
3080		3085		3090
Ile Leu	His Lys Leu Asp	Gln Leu Asp Ala Ser Arg	Gly Arg Ala	
3095		3100		3105
Ile Pro	Phe Cys Gly Gln	Arg Gly Arg Phe Lys Tyr	Glu Ile Leu	
3110		3115		3120
Val Lys	Thr Gly Trp Gly	Arg Gly Ser Gly Thr Thr	Ala His Val	
3125		3130		3135
Gly Ile	Met Leu Tyr Gly	Val Asp Ser Arg Ser Gly	His Arg His	
3140		3145		3150



Leu Asp 3155	Gly Asp Arg Ala Phe 3160	His Arg Asn Ser Leu Asp Ile Phe 3165
Arg Ile 3170	Ala Thr Pro His Ser 3175	Leu Gly Ser Val Trp Lys Ile Arg 3180
Val Trp 3185	His Asp Asn Lys Gly 3190	Leu Ser Pro Ala Trp Phe Leu Gln 3195
His Val 3200	Ile Val Arg Asp Leu 3205	Gln Thr Ala Arg Ser Ala Phe Phe 3210
Leu Val 3215	Asn Asp Trp Leu Ser 3220	Val Glu Thr Glu Ala Asn Gly Gly 3225
Leu Val 3230	Glu Lys Glu Val Leu 3235	Ala Ala Ser Asp Ala Ala Leu Leu 3240
Arg Phe 3245	Arg Arg Leu Leu Val 3250	Ala Glu Leu Gln Arg Gly Phe Phe 3255
Asp Lys 3260	His Ile Trp Leu Ser 3265	Ile Trp Asp Arg Pro Pro Arg Ser 3270
Arg Phe 3275	Thr Arg Ile Gln Arg 3280	Ala Thr Cys Cys Val Leu Leu Ile 3285
Cys Leu 3290	Phe Leu Gly Ala Asn 3295	Ala Val Trp Tyr Gly Ala Val Gly 3300
Asp Ser 3305	Ala Tyr Ser Thr Gly 3310	His Val Ser Arg Leu Ser Pro Leu 3315
Ser Val 3320	Asp Thr Val Ala Val 3325	Gly Leu Val Ser Ser Val Val Val 3330
Tyr Pro 3335	Val Tyr Leu Ala Ile 3340	Leu Phe Leu Phe Arg Met Ser Arg 3345
Ser Lys 3350	Val Ala Gly Ser Pro 3355	Ser Pro Thr Pro Ala Gly Gln Gln 3360
Val Leu 3365	Asp Ile Asp Ser Cys 3370	Leu Asp Ser Ser Val Leu Asp Ser 3375
Ser Phe 3380	Leu Thr Phe Ser Gly 3385	Leu His Ala Glu Gln Ala Phe Val 3390
Gly Gln 3395	Met Lys Ser Asp Leu 3400	Phe Leu Asp Asp Ser Lys Ser Leu 3405
Val Cys 3410	Trp Pro Ser Gly Glu 3415	Gly Thr Leu Ser Trp Pro Asp Leu 3420
Leu Ser 3425	Asp Pro Ser Ile Val 3430	Gly Ser Asn Leu Arg Gln Leu Ala 3435
Arg Gly 3440	Gln Ala Gly His Gly 3445	Leu Gly Pro Glu Glu Asp Gly Phe 3450

Ser	Leu	Ala	Ser	Pro	Tyr	Ser	Pro	Ala	Lys	Ser	Phe	Ser	Ala	Ser
3455						3460					3465			
Asp	Glu	Asp	Leu	Ile	Gln	Gln	Val	Leu	Ala	Glu	Gly	Val	Ser	Ser
3470						3475					3480			
Pro	Ala	Pro	Thr	Gln	Asp	Thr	His	Met	Glu	Thr	Asp	Leu	Leu	Ser
3485						3490					3495			
Ser	Leu	Ser	Ser	Thr	Pro	Gly	Glu	Lys	Thr	Glu	Thr	Leu	Ala	Leu
3500						3505					3510			
Gln	Arg	Leu	Gly	Glu	Leu	Gly	Pro	Pro	Ser	Pro	Gly	Leu	Asn	Trp
3515						3520					3525			
Glu	Gln	Pro	Gln	Ala	Ala	Arg	Leu	Ser	Arg	Thr	Gly	Leu	Val	Glu
3530						3535					3540			
Gly	Leu	Arg	Lys	Arg	Leu	Leu	Pro	Ala	Trp	Cys	Ala	Ser	Leu	Ala
3545						3550					3555			
His	Gly	Leu	Ser	Leu	Leu	Leu	Val	Ala	Val	Ala	Val	Ala	Val	Ser
3560						3565					3570			
Gly	Trp	Val	Gly	Ala	Ser	Phe	Pro	Pro	Gly	Val	Ser	Val	Ala	Trp
3575						3580					3585			
Leu	Leu	Ser	Ser	Ser	Ala	Ser	Phe	Leu	Ala	Ser	Phe	Leu	Gly	Trp
3590						3595					3600			
Glu	Pro	Leu	Lys	Val	Leu	Leu	Glu	Ala	Leu	Tyr	Phe	Ser	Leu	Val
3605						3610					3615			
Ala	Lys	Arg	Leu	His	Pro	Asp	Glu	Asp	Asp	Thr	Leu	Val	Glu	Ser
3620						3625					3630			
Pro	Ala	Val	Thr	Pro	Val	Ser	Ala	Arg	Val	Pro	Arg	Val	Arg	Pro
3635						3640					3645			
Pro	His	Gly	Phe	Ala	Leu	Phe	Leu	Ala	Lys	Glu	Glu	Ala	Arg	Lys
3650						3655					3660			
Val	Lys	Arg	Leu	His	Gly	Met	Leu	Arg	Ser	Leu	Leu	Val	Tyr	Met
3665						3670					3675			
Leu	Phe	Leu	Leu	Val	Thr	Leu	Leu	Ala	Ser	Tyr	Gly	Asp	Ala	Ser
3680						3685					3690			
Cys	His	Gly	His	Ala	Tyr	Arg	Leu	Gln	Ser	Ala	Ile	Lys	Gln	Glu
3695						3700					3705			
Leu	His	Ser	Arg	Ala	Phe	Leu	Ala	Ile	Thr	Arg	Ser	Glu	Glu	Leu
3710						3715					3720			
Trp	Pro	Trp	Met	Ala	His	Val	Leu	Leu	Pro	Tyr	Val	His	Gly	Asn
3725						3730					3735			
Gln	Ser	Ser	Pro	Glu	Leu	Gly	Pro	Pro	Arg	Leu	Arg	Gln	Val	Arg
3740						3745					3750			
Leu	Gln	Glu	Ala	Leu	Tyr	Pro	Asp	Pro	Pro	Gly	Pro	Arg	Val	His

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Thr Cys Ser Ala Ala Gly	Gly Phe Ser Thr Ser	Asp Tyr Asp Val		
3770	3775	3780		
Gly Trp Glu Ser Pro His	Asn Gly Ser Gly Thr	Trp Ala Tyr Ser		
3785	3790	3795		
Ala Pro Asp Leu Leu Gly	Ala Trp Ser Trp Gly	Ser Cys Ala Val		
3800	3805	3810		
Tyr Asp Ser Gly Gly Tyr	Val Gln Glu Leu Gly	Leu Ser Leu Glu		
3815	3820	3825		
Glu Ser Arg Asp Arg Leu	Arg Phe Leu Gln Leu	His Asn Trp Leu		
3830	3835	3840		
Asp Asn Arg Ser Arg Ala	Val Phe Leu Glu Leu	Thr Arg Tyr Ser		
3845	3850	3855		
Pro Ala Val Gly Leu His	Ala Ala Val Thr Leu	Arg Leu Glu Phe		
3860	3865	3870		
Pro Ala Ala Gly Arg Ala	Leu Ala Ala Leu Ser	Val Arg Pro Phe		
3875	3880	3885		
Ala Leu Arg Arg Leu Ser	Ala Gly Leu Ser Leu	Pro Leu Leu Thr		
3890	3895	3900		
Ser Val Cys Leu Leu Leu	Phe Ala Val His Phe	Ala Val Ala Glu		
3905	3910	3915		
Ala Arg Thr Trp His Arg	Glu Gly Arg Trp Arg	Val Leu Arg Leu		
3920	3925	3930		
Gly Ala Trp Ala Arg Trp	Leu Leu Val Ala Leu	Thr Ala Ala Thr		
3935	3940	3945		
Ala Leu Val Arg Leu Ala	Gln Leu Gly Ala Ala	Asp Arg Gln Trp		
3950	3955	3960		
Thr Arg Phe Val Arg Gly	Arg Pro Arg Arg Phe	Thr Ser Phe Asp		
3965	3970	3975		
Gln Val Ala His Val Ser	Ser Ala Ala Arg Gly	Leu Ala Ala Ser		
3980	3985	3990		
Leu Leu Phe Leu Leu Leu	Val Lys Ala Ala Gln	His Val Arg Phe		
3995	4000	4005		
Val Arg Gln Trp Ser Val	Phe Gly Lys Thr Leu	Cys Arg Ala Leu		
4010	4015	4020		
Pro Glu Leu Leu Gly Val	Thr Leu Gly Leu Val	Val Leu Gly Val		
4025	4030	4035		
Ala Tyr Ala Gln Leu Ala	Ile Leu Leu Val Ser	Ser Cys Val Asp		
4040	4045	4050		
Ser Leu Trp Ser Val Ala	Gln Ala Leu Leu Val	Leu Cys Pro Gly		
4055	4060	4065		

Thr Gly Leu Ser Thr Leu Cys Pro Ala Glu Ser Trp His Leu Ser  
 4070 4075 4080  
 Pro Leu Leu Cys Val Gly Leu Trp Ala Leu Arg Leu Trp Gly Ala  
 4085 4090 4095  
 Leu Arg Leu Gly Ala Val Ile Leu Arg Trp Arg Tyr His Ala Leu  
 4100 4105 4110  
 Arg Gly Glu Leu Tyr Arg Pro Ala Trp Glu Pro Gln Asp Tyr Glu  
 4115 4120 4125  
 Met Val Glu Leu Phe Leu Arg Arg Leu Arg Leu Trp Met Gly Leu  
 4130 4135 4140  
 Ser Lys Val Lys Glu Phe Arg His Lys Val Arg Phe Glu Gly Met  
 4145 4150 4155  
 Glu Pro Leu Pro Ser Arg Ser Ser Arg Gly Ser Lys Val Ser Pro  
 4160 4165 4170  
 Asp Val Pro Pro Pro Ser Ala Gly Ser Asp Ala Ser His Pro Ser  
 4175 4180 4185  
 Thr Ser Ser Ser Gln Leu Asp Gly Leu Ser Val Ser Leu Gly Arg  
 4190 4195 4200  
 Leu Gly Thr Arg Cys Glu Pro Glu Pro Ser Arg Leu Gln Ala Val  
 4205 4210 4215  
 Phe Glu Ala Leu Leu Thr Gln Phe Asp Arg Leu Asn Gln Ala Thr  
 4220 4225 4230  
 Glu Asp Val Tyr Gln Leu Glu Gln Gln Leu His Ser Leu Gln Gly  
 4235 4240 4245  
 Arg Arg Ser Ser Arg Ala Pro Ala Gly Ser Ser Arg Gly Pro Ser  
 4250 4255 4260  
 Pro Gly Leu Arg Pro Ala Leu Pro Ser Arg Leu Ala Arg Ala Ser  
 4265 4270 4275  
 Arg Gly Val Asp Leu Ala Thr Gly Pro Ser Arg Thr Pro Leu Arg  
 4280 4285 4290  
 Ala Lys Asn Lys Val His Pro Ser Ser Thr  
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 ccattcacct gctgtgtgac ctggtaaat

<210> 4  
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<223> PCR primer BPR9

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26

<210> 5  
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31

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21

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27

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<400> 9 20  
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<210> 11  
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<400> 11 21  
 agcgcaacta cttggaggcc c

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<210> 13  
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<400> 13 26  
 gaggtgtgg ggggccagtc aagtgg

<210> 14  
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 agggaggcag aggaaagggc cgaac

25

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<400> 15  
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24

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<400> 16  
 aagcgcaaaa gggctgcgctc g

21

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 ggccctccct gccttctagg cg

22

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<400> 18  
 gttgcagcca agcccatggtt a

21

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<400> 19

ggtcgcgctg tggcgaagg

19

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cggcgggcgg catcgt

16

<210> 21

<211> 16

<212> DNA

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<223> PCR primer 1F2

<400> 21

acggcggggc catgcg

16

<210> 22

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<212> DNA

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<223> PCR primer 1R2

<400> 22

gcgtcctggc ccgcgtcc

18

<210> 23

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<212> DNA

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<223> PCR primer 2F

<400> 23

ttggggatgc tggcaatgtg

20

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<210> 30  
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<400> 31  
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<400> 33  
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<210> 34  
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<223> PCR primer 5R3

<400> 34  
gaaggggagt gggcagcaga c

21

<210> 35  
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<400> 35  
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21

<210> 36  
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<400> 36  
tgccccagtg cttcagagat c

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<210> 37  
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<400> 37  
ggagtgccct gagccccct

19

<210> 38  
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<400> 38  
cccctaacca cagccagcg

19

<210> 39  
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<400> 39  
tctgttcgtc ctggtgtcct g

21

<210> 40  
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<400> 40  
gcaggagggc aggttgtaga a

21

<210> 41  
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<400> 41  
ggtaggggga gtctgggctt

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<210> 42  
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<400> 42  
gaggccaccc cgagtcc

17

<210> 43  
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<400> 43  
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<210> 44  
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<220>  
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<400> 44  
ggaaggtggc ctgaggagat 20

<210> 45  
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<400> 45  
ggggtccacg ggccatg 17

<210> 46  
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<400> 46  
aagcccagca gcacggtgag 20

<210> 47  
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<223> PCR primer 11midF

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gcttgacagcc acggaac 17

<210> 48  
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<220>  
<223> PCR primer 11midR

<400> 48  
gcagtgtac cactgagaac 20

<210> 49  
<211> 23  
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<400> 49

tgcccctggg agaccaacga tac

23

<210> 50  
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 <212> DNA  
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<220>  
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<400> 50  
 ggctgctgcc ctcaactggga ag

22

<210> 51  
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<220>  
 <223> PCR primer 12F

<400> 51  
 gaggcgacag gctaaggg

18

<210> 52  
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<220>  
 <223> Primer for PCR

<400> 52  
 aggtcaacgt gggcctccaa gtagt

25

<210> 53  
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<220>  
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<400> 53  
 gccttgcgca gcttggact

19

<210> 54  
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<220>  
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20

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30

<210> 56  
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Trp Asp Phe Gly Asp Gly Ser  
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<210> 59  
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His Leu Thr Ala  
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gcagggtgag caggtggggc catccta

27

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<213> Artificial sequence

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